

logo

NTDB id 1003164 ABHI22 RS12600 WP 003488165.1	.MARGINKVILVGNLGNDDPKYTQAGMAITRVSLATTSMRKDR.DGNNQERTEWHRVVFEGKLGEIAGEYLRKGSQVYV	78
NTDB id 1103 NMB RS07590 WP 002212976.1	...MSLNKVLIGRLGRDPEVRYMPNGEAVCNFVSATSETWNR.NGQRVTEWHRNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1131 NGFG RS05740 WP 003695064.1	...MSLNKVLIGRLGRDPEVRYMPNGEAVCNFVSATSETWNR.NGQRVTEWHRNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1166 A1552VC RS00795 WP 000168289.1	MASRGVNKVLIGNLGDDPEVRYMPSSGAVANITITATSETWRDKATGEQKEKTEWHRVITLYGKLAEVAGEYLRKGSQVYI	80
NTDB id 1390 A4U84 RS00055 WP 021115958.1	..MAGVNKVIIVGNLGNDDPMRTMPNGDAVATLSVATSESWNDKMTGERREVTWHRIVFRRQAEVAGQYLRKGSKVYV	78
consensus	*!!!!!!*!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!	

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NTDB id 1003164 ABHI22 RS12600 WP 003488165.1	EGRLQTRKYQGGQERYSTDIVA...NEMQMLGGRGEGGGGMSDRP.QRSAPRQQGGGGGGGGGGGYGG.....	145
NTDB id 1103 NMB RS07590 WP 002212976.1	EGRIQSRKYQGGKDGIERAYDIVA...NEMKMLGGRNENSGGAPYEYGYGQSQE.....AYQRPAQQSRQP	139
NTDB id 1131 NGFG RS05740 WP 003695064.1	EGRIQSRKYQGGKDGIERAYDIVA...NEMKMLGGRNENSGGAPYDEGYGQSQE.....AYQRPAQQSRQP	139
NTDB id 1166 A1552VC RS00795 WP 000168289.1	EGQLQTRKWQDDQSGQDRYSTEVVVQGYNGIMQMLGGRAGQG.CMPAQGGM...NVPAQ.....QGSWGGPPQPAK..	146
NTDB id 1390 A4U84 RS00055 WP 021115958.1	EGKLKTRKWQDDQNGQDRYTTEIQG...DVLQMLDSRSSGG.DFGCNQGSQWNQAPAQNTNY...NQGGYSDNYAQNN..	147
consensus	!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!*!!!!!!	

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NTDB id 1003164 ABHI22 RS12600 WP 003488165.1	...GGGGQDYAPRRQPPAQQSAPPMDDFADDDIPF	178
NTDB id 1103 NMB RS07590 WP 002212976.1	ASDAPSHPPQEAPAPRRQPVPAAAPVEDI.DDDIPF	174
NTDB id 1131 NGFG RS05740 WP 003695064.1	APDAPSHPPQEAPAPRRQPVPAAAPVEDI.DDDIPF	174
NTDB id 1166 A1552VC RS00795 WP 000168289.1	..QHQ..PMQQSAPQQYSQPQYNEPPMDF.DDDIPF	177
NTDB id 1390 A4U84 RS00055 WP 021115958.1	..NFNGGNATRPQPAQKPAAQAEPMDNF.DDDIPF	180
consensus	* * *** * * *** !**** !!!!!!!	

- non conserved
- similar
- ≥ 50% conserved